Figure 1 Plasmid pCMV.Bx08.gp160

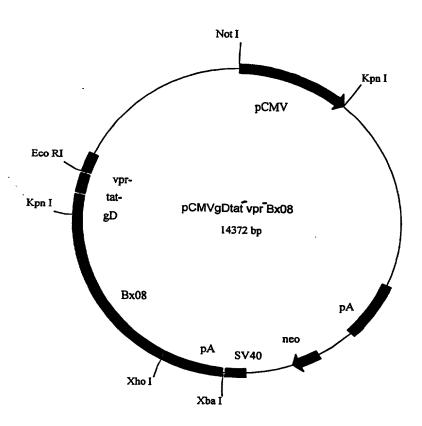


Figure 2 DNA immunization plasmid pCMV3Bx08.

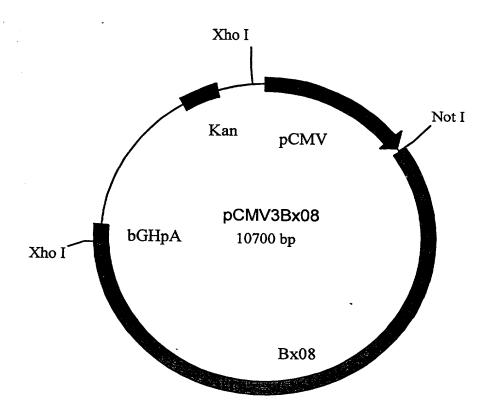


Figure 3. Pseudovirion Expression Plasmid p133B1 HIV-1 Bx08

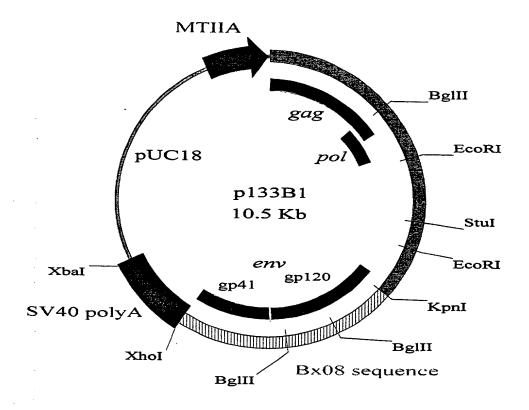
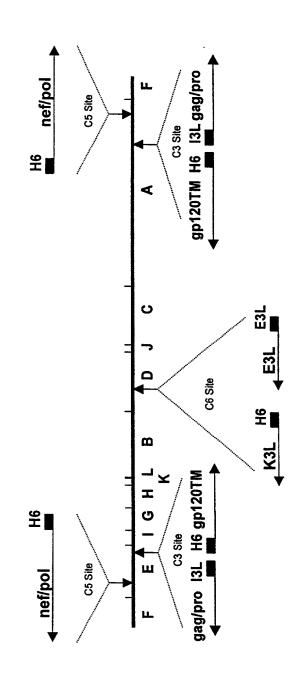
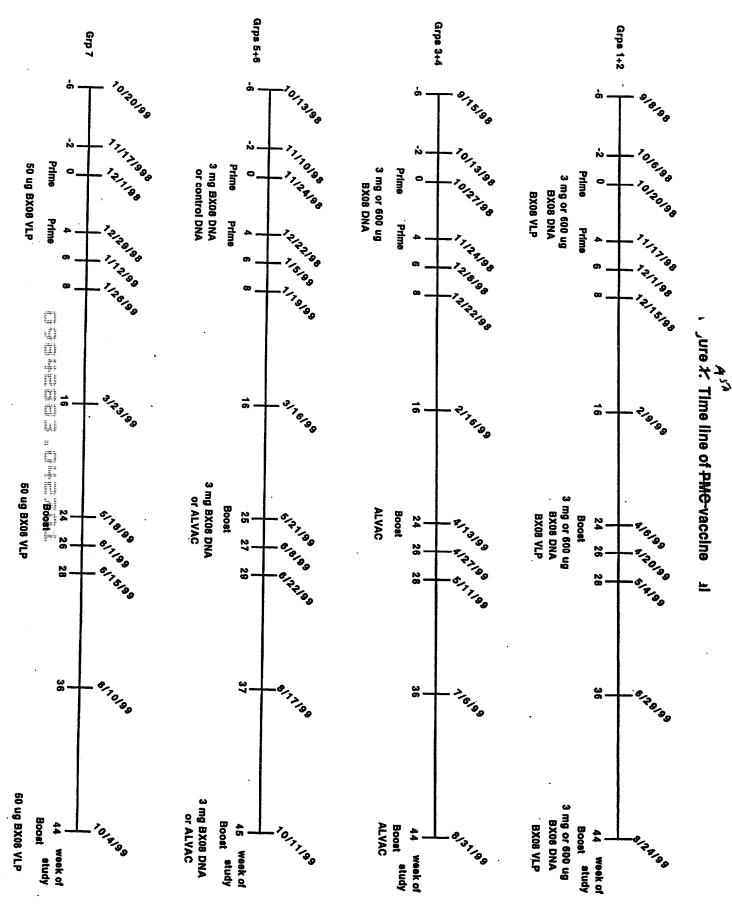


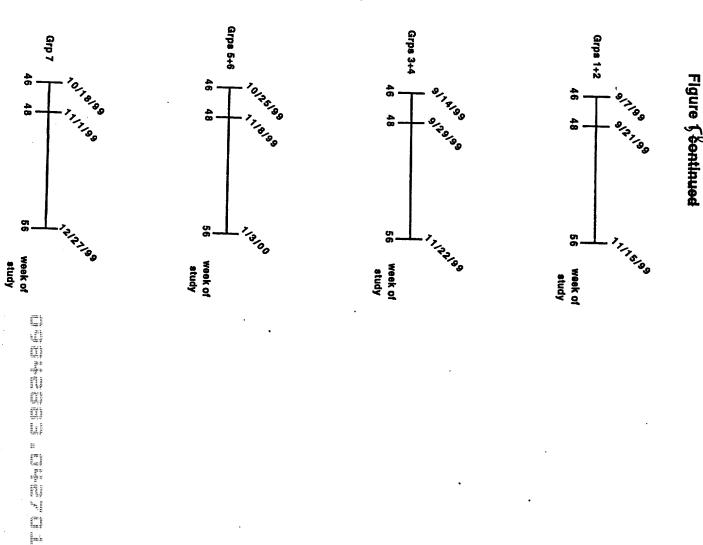
FIGURE 4

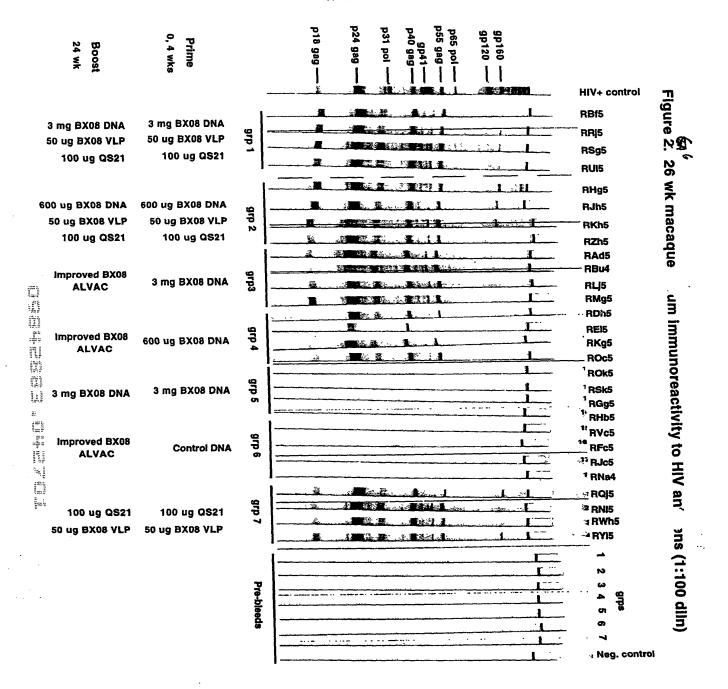
ALVAC(2)120(BX08)GNP (vCP1579)

(ALVAC Xhol Restriction Map)











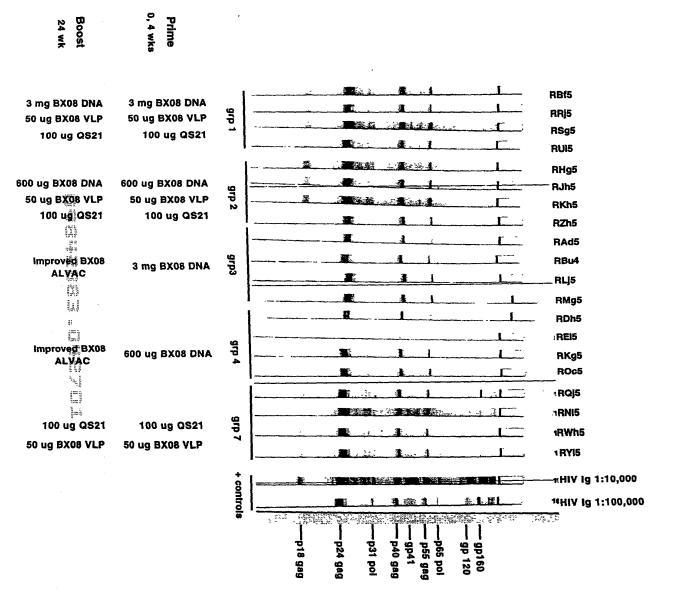


Figure 10 Plasmid pHIV76

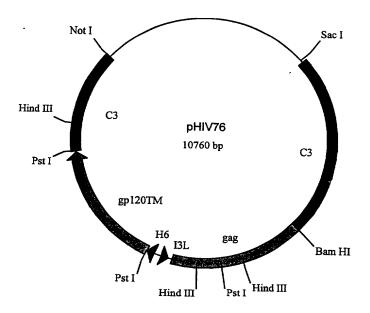


Figure 11 vCP1579: H6/HIV Pol/Nef epitope cassette in ALVAC C5 site

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1 TTTTTTCAT TATTTAGAAA TTATGCATTT TAGATCTTTA TAAGCGGCCG TGATTAACTA
  61 GTCATAAAAA CCCGGGATCG ATTCTAGACT CGAGGGTACC GGATCTTAAT TAATTAGTCA
 121 TCAGGCAGGG CGAGAACGAG ACTATCTGCT CGTTAATTAA TTAGGTCGAC GGATCCCCCA
 181 ACAAAACTA ATCAGCTATC GGGGTTAATT AATTAGTTAT TAGACAAGGT GAAAACGAAA
 241 CTATTTGTAG CTTAATTAAT TAGAGCTTCT TTATTCTATA CTTAAAAAGT GAAAATAAAT
 301 ACAAAGGTTC TTGAGGGTTG TGTTAAATTG AAAGCGAGAA ATAATCATAA ATTATTTCAT
 361 TATCGCGATA TCCGTTAAGT TTGTATCGTA ATGCCACTAA CAGAAGAAGC AGAGCTAGAA
 421 CTGGCAGAAA ACAGAGAGAT TCTAAAAGAA CCAGTACATG GAGTGTATTA TGACCCATCA
 481 AAAGACTTAA TAGCAGAAAT ACAGAAGCAG GGGCAAGGCC AATGGACATA TCAAATTTAT
 541 CAAGAGCCAT TTAAAAATCT GAAAACAGGA ATGGAGTGGA GATTTGATTC TAGATTAGCA
 601 TTTCATCACG TAGCTAGAGA ATTACATCCT GAATATTTTA AAAATTGTAT GGCAATATTC
 661 CAAAGTAGCA TGACAAAAAT CTTAGAGCCT TTTAGAAAAC AAAATCCAGA CATAGTTATC
 721 TATCAATACA TGGATGATTT GTATGTAGGA TCTGACTTAG AAATAGGGCA GCATAGAACA
 781 AAAATAGAGG AGCTGAGACA ACATCTGTTG AGGTGGGGAC TTACAACCAT GGTAGGTTTT
 841 CCAGTAACAC CTCAAGTACC TTTAAGACCA ATGACTTACA AAGCAGCTGT AGATCTTTCT
 901 CACTTTTAA AAGAAAAAGG AGGTTTAGAA GGGCTAATTC ATTCTCAACG AAGACAAGAT
 961 ATTCTTGATT TGTGGATTTA TCATACACAA GGATATTTTC CTGATTGGCA GAATTACACA
1021 CCAGGACCAG GAGTCAGATA CCCATTAACC TTTGGTTGGT GCTACAAGCT AGTACCAATG
1081 ATTGAGACTG TACCAGTAAA ATTAAAGCCA GGAATGGATG GCCCAAAAGT TAAACAATGG
1141 CCATTGACAG AAGAAAAAAT AAAAGCATTA GTAGAAATTT GTACAGAGAT GGAAAAGGAA
1201 GGGAAAATTT CAAAAATTGG GCCTTAATTT TTCTGCAGCC CGGGGGATCC TTTTTATAGC
1261 TAATTAGTCA CGTACCTTTG AGAGTACCAC TTCAGCTACC TCTTTTGTGT CTCAGAGTAA
1321 CTTTCTTTAA TCAATTCCAA AACAG
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Upstream (right) flanking sequence: 1-266

VV H6 promoter: 267-390

HIV pol/nef/pol/nef/pol cassette: 391-1227

Downstream (left) flanking sequence: 1227-1345

Figure 12 E3L and K3L genes in C6

10 20		* *	* *	* *		* *		* *
GAGCTCGCGG CCGCCTATCA A CTCGAGCGCC GGCGGATAGT 1	AAAGTCTTAA TGAGTTAGGT TTTCAGAATT ACTCAATCCA	GTAGATAGTA : CATCTATCAT I	TAGATATTAC ATCTATAATG	TACAAAGGTA ATGTTTCCAT	TTCATATITC AAGTATAAAG	CTATCAATTC GATAGTTAAG	TAAAGTAGAT ATTTCATCTA	GATATTAATA CTATAATTAT
120 130			170					
ACTCAAAGAT GATGATAGTA C	GATAATAGAT ACGCTCATAT	AATGACTGCA	AATTTGGACG	GTTCACATTT	TAATCATCAC	GCGTTCATAA	GTTTCAACTG	CATAGATCAA
230 240	250 260	270	280	290	300	310	320	330
AATCTCACTA AAAAGATAGC C	CGATGTATTT GAGAGAGATT	GGACATCTAA	CTACGCTAAA	GAAATTACAG	AATAAATAA	TACATAATGG	ATTTTGTTAT	CATCAGITAT
indicioni illicinico	GUINGHIANA CICICICIA	. CCIGIAGAII	GATGCGATT	CITIAAIGIC	AALAITIAIT	AIGIATIACC	TAAAACAATA	GIAGICAATA
340 350 * * * *	* * *		* *			* *	430	* *
ATTTAACATA AGTACAATAA A TAAATTGTAT TCATGTTATT 1	AAAGTATTAA ATAAAAATAC TITCATAATT TATITITATG	TTACTTACGA A	AAAAATGACT TTTTTACTGA	AATTAGCTAT TTAATCGATA	AAAAACCCAG TTTTTGGGTC	ATCTCTCGAG TAGAGAGCTC	GTCGACGGTA CAGCTGCCAT	TCGATAAGCT AGCTATTCGA
450 460 * * * *	470	480	490	500		.0	520	530
TGATATCGAA TTCATAAAAA T ACTATAGCTT AAGTATTTTT A	TT A TTG ATG TCT ACA AA T AAC TAC AGA TGT <q c<="" h="" r="" td=""><td>GTA GGA AAA (M R K</td><td>CAT TAA CTC Y N V</td><td>TAG ATA TA</td><td>TA TCC TTT T AT AGG AAA I 'G K</td><td>GT ATA ATC CA TAT TAG T Y D</td><td>AAC TCT AAT TTG AGA TTI V R I</td><td>V K</td></q>	GTA GGA AAA (M R K	CAT TAA CTC Y N V	TAG ATA TA	TA TCC TTT T AT AGG AAA I 'G K	GT ATA ATC CA TAT TAG T Y D	AAC TCT AAT TTG AGA TTI V R I	V K
540 550	0 560			-			610	620
AAC TIT TAC AGT TIT CCC	* * * *	ATA TTC AAC	* * * *	בית אלים שלים שלים	* יאם דידי אאר	* * * **	* 1	
TTG AAA ATG TCA AAA GGG	g atg gtc aaa tag gga	TAT AAG TTG Y E V	TAT AGA TA	G GTA TAC (TA GAA TTG M K V	TGA GAG ACC	GTT CTA TO	CG AAG TCT A E S
630 640	0 650	660	670	686		590 * *	700	710
GTG AGG ATA GTC AAA AAG CAC TCC TAT CAG TTT TTC <h d="" f="" l<="" p="" td="" y=""><td>G ATA AAT GTA TAG AGO C TAT TTA CAT ATC TCG Y I Y L A</td><td>ATA ATC CTT TAT TAG GAA Y D K</td><td>CTC GTA TA GAG CAT AT E Y V</td><td>C TCT GCC (G AGA CGG (R G</td><td>TT TAT TAC AAA ATA ATG K I V</td><td>TAG CGG GCC</td><td>TAA CCC G</td><td>AA CGA ATA FT GCT TAT L S Y</td></h>	G ATA AAT GTA TAG AGO C TAT TTA CAT ATC TCG Y I Y L A	ATA ATC CTT TAT TAG GAA Y D K	CTC GTA TA GAG CAT AT E Y V	C TCT GCC (G AGA CGG (R G	TT TAT TAC AAA ATA ATG K I V	TAG CGG GCC	TAA CCC G	AA CGA ATA FT GCT TAT L S Y
				КЗЪ				
720 730	0 740 7	50 76	0 77	0 75	10 71	90 80	ın 8 [.]	10
ACA AAA TGC AAG CAT ACG TGT TIT ACG TTC GTA TGC <c a="" f="" l="" m<="" td=""><td>* * * * * G ATACAAACTT AACGGATA</td><td>50 76 * * TC GCGATAATG. AG CGCTATTAC</td><td>* * A AATAATTI</td><td>* * AT GATTATITO</td><td>T CGCTTTCAI</td><td>* * AT TTAACACAI</td><td>0 8: * * AC CCTCAAGA FG GGAGTTCT</td><td>AC</td></c>	* * * * * G ATACAAACTT AACGGATA	50 76 * * TC GCGATAATG. AG CGCTATTAC	* * A AATAATTI	* * AT GATTATITO	T CGCTTTCAI	* * AT TTAACACAI	0 8: * * AC CCTCAAGA FG GGAGTTCT	AC
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC C F A L M	G ATACAAACTT AACGGATA C TATGTTTGAA TTGCCTAT	* TC GCGATAATG AG CGCTATTAC	* * A AATAATTII T TTATTAAAI	* * AT GATTATTT TA CTAATAAA	* CGCTTTCA: EA GCGAAAGT	* TTAACACAI	C CCTCAAGA	AC TG
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC C F A L M	ATACAAACIT AACGGATA C TATGITIGAA TIGCCIAI 840 850	TC GCGATAATG AG CGCTATTAC 860	* AATAATITA T TTATTAAA 870	* CAGATTGTTT	T CGCTTTCAL EA GCGAAAGT 890 CGTTTTCCCC	TTGGCGTATC	C CCTCAAGA G GGAGTTCT	920
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M	* ATACARACTT AACGGATA C TATGTTIGAA TIGCCTAT 840 850 CTTAAGTATA GAATAAAGAA AAATTCATAT CITATTCTI	TC GCGATAATG AG CGCTATTAC 860 * AGCTCTAATT TCGAGATTAA	* A AATAATITI T TTATTAAA 870 * AATTAATGAA TTAATTACTT	* T GATTATITY CA CTAATAAM 880 * CAGATTGTTT GTCTAACAAA	* CGTTTCAL 890 * CGTTTCCCC GCARAGGGG	TIGGGGTATC AACCGCATAG	C CCTCAAGA C GGAGTTCT 910 * ACTAATTAAT TGATTAATTA	P20 TAACCCGGGC
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M 820 830 CTTTGTATTT ATTTCACTT T GAAACATAAA TAAAAGTGAA A	840 850 TTTAAGTATA GAATAAAGAAAAATTCATAT CTTATTCTT	* CGCATAATG AG CGCTATTAC 860 * AGCTCTAATT TCGAGATTAA	* A AATAATITI T TTATTAAAT 870 AATTAATGAA TTAATTACTT 980	* * IT GATTATITY TA CTAATAAM 880 * * CAGATTGTTT GTCTAACAAA 990 * *	* CGCTTCA: 890 CGTTTCCCC GCAAAAGGGG	TTGCGTATC AACCGCATAG	C CCTCAAGA: G GGAGTTCT 910 ACTAATTAAT TGATTAATTA 1020	P20 TAACCCGGGC
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M S20 830 CTTTGTATTT ATTTTCACTT T GAAACATAAA TAAAAGTGAA A	S ATACAAACTT AACGGATA C TATGTITGAA TIGCCTAT 840 850 ITTAAGTATA GAATAAAGAA AAATTCATAT CITAITTCTI 950 960	* CGCATAATG AG CGCTATTAC 860 * AGCTCTAATT * TCGAGATTAA 970 * GTATACACAT	* A AATAATTII T TTATTAAAT 870 * AATTAATGAA TTAATTACIT 980 * AACCATTACT	* T GATTATITY CA CTAATAAN 880 * CAGATTGTTT GTCTAACAAA 990 * AACGTAGAAT	* CGCTTCA: A GCGAAAGT 890 * CGTTTCCCC GCAAAAGGGG 1000 * TATAGGAAG	TTGGCGTATC AACCGCATAG AACCGCATAG AACCGCATAG AACCGCATAG AACCGCATAG	GGAACAGGGT	P20 TAACCCGGGC ATTGGGCCCG
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M 820 830 CTTTGTATTT ATTTTCACTT T GAAACATAAA TAAAAGTGAA A 930 940 TGCAGCTCGA GGAATTCAAC T	S ATACAAACTT AACGGATA C TATGTITGAA TIGCCTAT 840 850 ITTAAGTATA GAATAAAGAA AAATTCATAT CITAITTCTI 950 960	* CGCATATAC 860 * AGCTCTAATT * TCGAGATTAA 970 * GTATACACAT * CATATGTGTA	* A AATAATTTA T TTATTAAAT 870 ** AATTAATGAA TTAATTACTT 980 ** AACCATTACT TTGGTAATGA 1090	* * TGATTATTIC *A CTAATAAAC ** 880 ** CAGATTGTTT GTCTAACAAA ** 990 ** AACGTAGAAT TTGCATCTTA	* T CGCTTTCA: R 900 **CGTTTTCCCC GCAAAAGGGG 1000 GTATAGGAAG CATATCCTTC	* * * * * * * * * * * * * * * * * * *	GGAACAGGGT	TAACCCGGGC ATTGGGCCCG 1030 TTGTTGATTC AACAACTAAG
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M 820 830 CTTTGTATTT ATTTTCACTT T GAAACATAAA TAAAAGTGAA A 930 940 TGCAGCTCGA GGAATTCAAC T ACGTCGAGCT CCTTAAGTTG A	840 850 840 850 FITTARGTATA GAATAAAGAA AAATTCATAT CTTATTCTT 950 960 FATATCGACA TATTCATT ATTATAGCTGT ATAAAGTAAA 1060 1070	* TGCACGTAATG ** ** ** ** ** ** ** ** ** ** ** ** *	* A AATAATTTI T TTATTAAAT 870 AATTAATGAA TTAATTACTT 980 AACCATTACT TTGGTAATGA 1090 ** CTATTATAGA	* TGCCAGATA	TCGATTTCAL 890 CGTTTTCCCC GCAAAAGGGG 1000 GTATAGGAAG CATATCCTTC 1110 TCTATATAAT	TTGGCGTATC AACCGCATAG 1010 AGAGTAACAG TCTACATTGC 1120 TATTTTGTAA	910 ACTAATTAAT TGATTAATTA 1020 GGAACAGGGT CCTTGTCCCA	P20 TAACCCGGGC ATTGGCCCG 1030 TTGTTGATTC AACAACTAAG 1140 ACTATGTGAT
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M 820 830 CTTTGTATTT ATTTTCACTT T GAAACATAAA TAAAAGTGAA A 930 940 TGCAGGTCGA GGAATTCAAC T ACGTCGAGCT CCTTAAGTTG A 1040 1050 GCAAACTATT CTARTACATA A	S ATACARACTT AACGGATE C TATGTTIGAN TIGCCTAT 840 850 FITTANGTATA GAATAAAGAE ANAITCATAT CTTATTCTT 950 960 FATATOGACA TATTCATTT ATATAGCAGT ATAAAGTAE 1060 1070 ATTCTTCTGT TAATACGTCT TAAAGAAGACA ATTATGCAGE 1170 1180	* TGCACGTAATG ** ** ** ** ** ** ** ** **	* A AATAATTI T TIATTAAAT 870 * AATTAATGAA TIAATTACT 980 * AACCATTACT TTGGTAATGA 1090 * CTATTATAGA GATAATATCT	* * * * * * * * * * * * * * * * * * *	* T CGCTTTCA: ** A GCGAAAGT ** 890 ** CGTTTTCCCC GCAAAAGGGG ** 1000 ** GTATAGGAAG CATATCCTTC ** 1110 ** TCTATATAAT AGATATATTA 1220	POOL TANACACATA POOL TIGGCGTATC AACCGCATAG AGATGTAACG TOTACATTGC TATITIGTAA ATAAAACATT 1230	910 ACTAATTAAT TGATTAATTA 1020 GGAACAGGGT CCTTGTCCCA 1130 GATGATGTA GATGATGTA 1240	TAACCCGGGC ATTGGGCCCG 1030 TIGTTGATTC AACAACTAAG 1140 ACTATGGAT TGATACACTA
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M 820 830 CTTTGTATTT ATTTCACTT T GAAACATAAA TAAAAGTGAA A 930 940 TGCAGCTCGA GGAATTCAAC T ACGTCGAGCT CCTTAAGTTG A 1040 1050 GCAAACTATT CTAATACATA A CGTTTGATAA GATTATGTAT T	840 850 840 850 FITTANGTATA GAATAAAGAA RAATTCATAT CTTATTCTT 950 960 FATATCGACA TATTCATT ATATAGCTGT ATAAAGTAAA ATATCCTAT TATTCATT 1060 1070 ATTCTTCTGT TAATACGTCT FAAGAAGACA ATTATGCAGA 1170 1180	* TTCCAACTTA	* A AATAATTI T TTATTAAAT 870 ** ** ** ** ** ** ** ** ** ** ** ** *	* * * * * * * * * * * * * * * * * * *	* T CGCTTTCA: ** S90 ** S90 ** CGTTTTCCCC ** GCAAAAGGGG ** 1000 ** GTATAGGAAG CATATCCTTC ** 1110 ** TCTATATAAT ** AGATATATTA ** 1220 ** CGTAATATCT	900 TTGGCGTATC AACCGCATAG 1010 AGATGTAACG TCTACATTGC 1120 TATTTTGTAA ATAAAACATT 1230 ATAGCATCCT	910 ACTAATTAAT TGATTAATTA 1020 GGAACAGGGT CCTTGTCCCA 1130 GATGATGTAA CTACTACAAT	P20 TAACCCGGGC ATTGGGCCCG 1030 TTGTTGATTC AACAACTAAG 1140 ACTATGTGAT TGATACACTA 1250 TATTCGCATAT
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M 820 830 CTTTGTATTT ATTTTCACTT T GAAACATAAA TAAAAGTGAA A 930 940 TGCAGGCTCGA GGAATTCAAC T ACGTCGAGCT CCTTAAGTTG A 1040 1050 GCAAACTATA CTAATACATA A CGTTTGATAA GATTATGTAT T 1150 1160 CTATATAAAGT AGTGTAATAA T	840 850 840 850 *** *** *** *** *** *** ***	* TC GCGATAATG AG CGCTATTAC ** ** ** ** ** ** ** ** ** ** ** ** *	* A AATAATTTI T TTATTAAAT 870 ** ** ** ** ** ** ** ** ** ** ** ** *	* * * * * * * * * * * * * * * * * * *	* T CGCTTTCA: ** 890 ** * CGTTTTCCCC ** GCAAAAGGGG ** 1000 ** GTATAGGAAG CATATCCTTC ** 1110 ** TCTATATAAT ** AGATATATTA ** 1220 ** CGTAATATCT GCATTATAGA 1330	900 TTGGCGTATC AACCGCATAG 1010 AGATGTAACG TCTACATTGC 1120 TATTTTGTAA ATAAAACATT 1230 ATAGCATCCT TATCGTAGGA	910 ACTAATTAAT TGATTAATTA 1020 GGAACAGGGT CCTTGTCCCA 1130 GATGATGTTA 1240 CAAAAAATAT GTTTTTTATA	P20 TAACCCGGGC ATTGGGCCCG 1030 TTGTTGATTC AACAACTAAG 1140 ACTATGTGAT TGATACACTA 1250 ATTCGCATAT TAAGCGTATA 1360
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M 820 830 CTTTGTATTT ATTTCACTT T GAAACATAAA TAAAAGTGAA A 930 940 TGCAGGCTCGA GGAATTCAAC T ACGTCGAGCT CCTTAAGTTG A 1040 1050 GCAAACTATT CTAATACATA A CGTTTGATAA GATTATGTAT T 1150 1160 CTATATAAGT AGTGTAATAA T GATATATTCA TCACATTAT A GATATATTCA TCACATTATT A	840 850 840 850 *** ** ** ** ** ** ** ** ** ** ** **	* * * * * * * * * * * * * * * * * * *	* A AATAATTIT T TTATTAAAT 870 AATTAATGAA TTAATTACT 980 AACCATTACT TTOGTAATGA 1090 CTATTATAGA GATAATACT 1200 GTCTTTGTGA CAGAAACACT 1310 ATAATAATCT	* * * * * * * * * * * * * * * * * * *	** TCTCATATATA	900 TTGGGGTATC AACCGCATAG 1010 AGATGTAACAG 1120 TATTTTGTAA ATAAAACATT 1230 ATAGCATCCT TATCGTAGGA 1340 ATTAATGATA	910 ACTAATTAAT 1020 GGAACAGGGT CCTTGTCCCA 1130 CAAAAAAATAT CTACTACAAT 1240 CAAAAAAATAT GTTTTTTTAA	P20 TAACCCGGGC ATTGGGCCCG 1030 TTGTTGATTC AACAACTAAG 1140 ACTATGTGAT TGATACACTA 1250 ATTCGCATAT TAAGCGTATA 1360 CACTATCTTC
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M 820 830 CTTTGTATTT ATTTTCACTT T GAAACATAAA TAAAAAGTGAA A 930 940 TGCAGCTCGA GGAATTCAAC T ACGTCGAGCT CCTTAAGTTG A 1040 1050 GCAAACTATT CTAATACATA A CGTTTGATAA GATTATGTAT T 1150 1160 CTATATAAGT AGGTTAATAA T GATATATCA TCACATTATT A 1260 1270 ATTCCCAAGT CTTCAGTTCT A TAAGGGTTCA GAAGTCAAGA T TAAGGGTTCA GAAGTCAAGA T 1370 1380	840 850 ETTARGTATA GATARAGAR ARATTCATAT 950 960 ETTARGTCATA TATATCGACA TATTCATT ATTATAGCATA ATTATAGATAR 1060 1070 ETTARGAGACA ATTATAGCAGA ATTCTTCTGT TARTAGCAGA 1170 1180 ETCATGTATA AGCTATATAGAGAGAAAATTCATATAAAGTAAAAATTCATATAAAAAAAA	* TTCCACCTTA 1190 * TTCCACCTTA 1190 * TTCCACCTTA 1190 * TTCCACCTTA 1300 * CGTATGGAAT 1310 * CGTATGGAAT 1410	* A AATAATTIT T TTATTAAAT * 870 * 870 * AATTAATGAA TTAATTACTT * 980 * AACCATTACT TTGGTAATGA 1090 * CTTATTATAGA GATAATACT 1200 * GTCTTATGGAACACT 1310 * ATAATAATCT TATTATTAGA	* * * * * * * * * * * * * * * * * * *	* TTCTGATATC GCATATATC CGTATATCATC 1110 CGTATAGAGAG CATATCCTTC 1110 TCTATATATA AGATATATTA 1220 CGTAATATCT GCATATATCT AGAGACATATC 1330 TTCTGATATC AAGACTATAG	900 TTGGCGTATC AACCGCATAG 1010 AGATGTAACG TCTACATTGC 1120 * TATTTTGTAA ATAAAACATT 1230 ATAGCATCCT TATCGTAGGA 1340 ATTAATGATA TAAATGATA	910 ACTAATTAAT 1020 GGAGCTCCCA 2130 GGAACAGGGT CCTTGTCCCA 2130 GAACAAGTTAATTA 1240 CAAAAAATTA 2240 CAAAAAATTA 1350 TAGTTTTTCA ATCAAAAACT	P20 TAACCCGGGC ATTGGCCCG 1030 TTGTTGATTC AACAACTAAG 1140 ACTATGTGAT TGATACACTA 1250 ATTCGCATAT TAAGCGTATA TAAGCGTATA CACTATCTTC GTGATACACAG
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CF A L M 820 830 CTTTGTATTT ATTTTCACTT T GAAACATAAA TAAAAGTGAA A 930 940 TGCAGGCTCGA GGAATTCAAC T ACGTCGAGCT CCTTAAGTTG A 1040 1050 GCAAACTATT CTAATACATA A CGTTTGATAA GATTATGTAT T 1150 1160 CTATATAAGT AGTGTAATAA T GATATATTCA TCACATTATT A 1260 1270 ATTCCCAAGT CTTCAGTTCT A TAAGGGTTCA GAAGTCAAGA T 1370 1380 TGTCAATTGA TTCTTATTCA C	840 850 TTTAAGTATA GAATAAAGAA AAATTCATAT 950 960 TATATTCTATATATATATATATATATATATATATATATA	* TC GCGATAATG 860 * 860 * AGCTCTAATT * TCGAGATTAA 970 * GTATACACAT CATATGTGTA 1080 * TGCACGTAAT ACGTGCATTA 1190 * TTCCAACTCT AAGGTTGAGA 1300 * CGTATACGAT CGTATACGAT 1410 * GCGTCCCTAG	* A AATAATTII T TTATTAAAT	* TGCCATTANT ** S80 ** S80 ** CAGATTGTTT GTCTAACAAA ** 990 ** AACGTAGAAT TTGCATCTTA ** 1100 ** TGCCAAGATA ACGGTTCTAT ** 1210 ** TGCCAAGATA ACAGATCAAA ** 1320 ** TGCCATTAAT ** TGCATTAAT ** TGCCATTAAT ** TGCCATTAAT ** TGCCATTAAT	* TTCTATATATA AGACTATAGA ** ** ** ** ** ** ** ** ** ** ** ** *	* TANACACAN ** 900 ** 900 ** TTGGCGTATC ** AACCGCATAG ** 1010 ** AGATGTAACG ** TCTACATTGC ** 1120 ** ** TATTTTGTAA ** ATTAAAACATT ** 1230 ** ATTAACATCAT ** 1340 ** ATTAATGATA ** TAATTACTAT ** TAGGTTCTGG	910 ACTAATTAAT 1020 GGAACAGGGT CCTTGTCCCA 1130 GAACAAATTAT 1240 CAAAAAATTA 1350 TAGTTTTTTGA ATCAAAAACT	P20 TAACCCGGGC ATTGGCCCG 1030 TTGTTGATTC AACAACTAAG 1140 ACTATGTGAT TGATACACTA 1250 ATTGGCATAT TAAGCGTATA TAAGCGTATA CACTATCTC GTGATACACA 1470 TCTATTATAC
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CC F A L M 820 830 CTTTGTATTT ATTTTCACTT T GAAACATAAA TAAAAGTGAA A 930 940 TGCAGGTCGA GGAATTCAAC T ACGTCGAGGT CCTTAAGTTG A 1040 1050 GCAAACTATT CTAATACATA A CGTTTGATAA GATTATGTAT T 1150 1160 CTATATAAAGT AGTGTAATAA T GATATATTCA TCACATTATT A 1260 1270 ATTCCCAAGT CTTCAGTTCT A TAAGGGTTCA GAAGTCAAGA T TAAGGGTTCA GAAGTCAAGA T	840 850 TTTAAGTATA GAATAAAGAA AAATTCATAT 950 960 TATATTCTATATATATATATATATATATATATATATATA	* TC GCGATAATG 860 * 860 * AGCTCTAATT * TCGAGATTAA 970 * GTATACACAT CATATGTGTA 1080 * TGCACGTAAT ACGTGCATTA 1190 * TTCCAACTCT AAGGTTGAGA 1300 * CGTATACGAT CGTATACGAT 1410 * GCGTCCCTAG	* A AATAATTII T TTATTAAAT	* TGCCATTANT ** S80 ** S80 ** CAGATTGTTT GTCTAACAAA ** 990 ** AACGTAGAAT TTGCATCTTA ** 1100 ** TGCCAAGATA ACGGTTCTAT ** 1210 ** TGCCAAGATA ACAGATCAAA ** 1320 ** TGCCATTAAT ** TGCATTAAT ** TGCCATTAAT ** TGCCATTAAT ** TGCCATTAAT	* TTCTATATATA AGACTATAGA ** ** ** ** ** ** ** ** ** ** ** ** *	* TANACACAN ** 900 ** 900 ** TTGGCGTATC ** AACCGCATAG ** 1010 ** AGATGTAACG ** TCTACATTGC ** 1120 ** ** TATTTTGTAA ** ATTAAAACATT ** 1230 ** ATTAACATCAT ** 1340 ** ATTAATGATA ** TAATTACTAT ** TAGGTTCTGG	910 ACTAATTAAT 1020 GGAACAGGGT CCTTGTCCCA 1130 GAACAAATTAT 1240 CAAAAAATTA 1350 TAGTTTTTTGA ATCAAAAACT	P20 TAACCCGGGC ATTGGCCCG 1030 TTGTTGATTC AACAACTAAG 1140 ACTATGTGAT TGATACACTA 1250 ATTGGCATAT TAAGCGTATA TAAGCGTATA CACTATCTC GTGATACACA 1470 TCTATTATAC
ACA AAA TGC AAG CAT ACG TGT TTT ACG TTC GTA TGC CF A L M 820 830 CTTTGTATTT ATTTTCACTT T GAAACATAAA TAAAAGTGAA A 930 940 TGCAGGCTCGA GGAATTCAAC T ACGTCGAGCT CCTTAAGTTG A 1040 1050 GCAAACTATT CTAATACATA A CGTTTGATAA GATTATGTAT T 1150 1160 CTATATAAGT AGTGTAATAA T GATATATTCA TCACATTATT A 1260 1270 ATTCCCAAGT CTTCAGTTCT A TAAGGGTTCA GAAGTCAAGA T 1370 1380 TGTCAATTGA TTCTTATTCA C	840 850 TTTAAGTATA GAATAAAGAA AAATTCATAT 950 960 TATATTCATAT ATATAGCATA TATATCATT ATATAGCATA TATATCATT ATATAGCATA TATATCATAT ATATAGCATA TATATCATAT ATATAGCATAT TCAATATACAGA 1170 1180 TTAATAGTATA TCAATATATCAGAA 1170 1180 TTAATATCATAT TCAATATATCAGAA 1280 1290 ATCTTCTAAA AAATCTCAATATATCAGAAGAATT TTTAGAAGTT TAGAAGATT TTTAGAAGTT 1390 1400 TTATATCTAAA GAAACGGATE SATATATCAAT GAAACGGATE SATATATCAAT CTTTGCCTAT	* TC GCGATAATG AG CGCTATTAC ** 860 ** ** ** ** AGCTCTAATT ** TCGAGATTAA ** 970 ** GTATACACAT ** CATATGTGTA ** 1080 ** TGCACGTAAT ** ACGTGCATTA ** 1190 ** TTCCAACTCT ** AAGGTTGAGA ** 1300 ** CGTATCGAAT ** GCATACCTTA ** 4110 ** 466GTCCCTAG ** CGCAGGGATC ** 1520 **	* A AATAATTII T TTATTAAAT	* * * * * * * * * * * * * * * * * * *	* TCGCTTTCAL ** 890 ** 890 ** CGTTTTCCCC GCAAAAGGGG ** 1000 GTATAGGAAG CATATCCTTC ** 1110 ** ** TCTATATAAT ** AGATATATT ** AGATATATTA ** 1220 ** CGTAATATATA ** 1330 ** TTCTGATATAC ** AAGACTATAGA ** AAGACTATAGA ** ATCTCTATTAT ** TAGAGATAAAT ** TAGAGATAAT ** TAGAG	* TANCACAN * 900 * 900 * TIGGCGTATC * AACTGGATAG * 1010 * AGATGTAACG * 1120 * * * * * * * * * * * * * * * * * * *	GAACAGGT CCTCAGAL GGAGTTCT 910 ACTAATTAAT 1020 GGAACAGGGT CCTTGTCCCA 1130 GATGATGTTA CTACTACAAT 1240 CAAAAAATAT 1350 TAGTTTTTGAT ATCAAAAACT 1460 ACATAATTCA TGTATTAAGT	P20 TAACCCGGGC ATTGGCCCG 1030 TTGTTGATTC AACAACTAAG 1140 ACTATGTGAT TGATACACTA 1250 ATTCGCATAT TAAGCGTATA CACTATCTCC GTGATAGAAG 1470 TCTATTATAC AGATAATATG

1610 1620 1630 1640 1650 1660 1670 ATAGTGACTA TITCATTCTC TGAAAATTGG TAACTCATTC TATATATGCT TTCCTTGTTG ATGAAGGATA GAATATACTC AATAGAATTT GTACCAACAA ACTGTTCTCT
TATCACTGAT AAAGTAAGAG ACTITTAACC ATTGAGTAAG ATATATACGA AAGGAACAAC TACTTCCTAT CTTATATGAG TTATCTTAAA CATGGTTGTT TGACAAGAGA 1760 1770 1710 1720 1730 1740 1750 TATGAATCGT ATATCATCAT CTGAAATAAT CATGTAAGGC ATACATTTAA CAATTAGAGA CTTGTCTCCT GITATCAATA TACTATTCTT GTGATAATTA TATGTGTGAGG ATACTTAGCA TATAGTAGTA GACTITATTA GTACATTCCG TATGTAAATT GTTAATCTCT GAACAGAGGA CAATAGTTAT ATGATAAGAA CACTATTAAA TACACACTCC 1810 1820 1830 1840 1850 1860 1870 1880 1890 1900 1910
CARATTEC CACCITCT ARTITICITA TACTAGATAT CARATCATA GAGCTACAG TICTIGGCT ARACAGATAT ACTITICIG GAACAATTC TACAACATTA GTITAAACAG GIGCAAGAA TIAAACAAT ATCATCATA GTITAGGTTA CCTCGATGT ARGAACCGAA TITGTCTATA TCARAAAGAC CTTGTTTAAG ATGTTGTAAT 1920 1930 1940 1950 1960 1970 1980 1990 2000 2010 2020
TTATABAGGA CTTTGGGTAG ATAAGTGGGA TGAAATCCCT ATTTAATTAA TGCTATCGCA TTGTCCTCGT GCAAATATCC AAACGCTTT GTGATAGTAT GGCATTCATT
AATATTTCCT GAAACCCATC TATTCACCCT ACTTTAGGAT AAAATTAATT ACGATAGCGT AACAGGAGCA CGTTTATAGG TTTGCGAAAA CACTATCATA CCGTAAGTAA 2520 2530 2540 2550 2560 2570 2780 2790 2800 2810 2820 2830 2840 2850 2860 2870 2880
CTAACACAC CAGCAATAAA ACTGAACCIA CTTIATCAT TITITATICA TCACCICTG GTGTTCGTC GTTCTATCG AATGTAGCT TGATTAACC GTCATCAATA
GATTGTGTTG GTCGTTATT TGACTTGGAT GAAATAGTA AAAAATAGT AGTAGGAGAC CACCAAGCAG CAAAGATAGG TTACATCGG ACTAATTGG CAGTAGATAT GGTGATGCTG GTTCTGGAGA TTCTGGAGGA GATGGATTAT TATCTGGAAG AATCTCTGTT ATTTCCTTGT TTTCATGTAT CGATTGCGTT GTAACATTAA GATTGCGAAA CCACTACGAC CAAGACCTCT AAGACCTCCT CTACCTAATA ATAGACCTTC TTAGAGACAA TAAAGGAACA AAAGTACATA GCTAACGCAA CATTGTAATT CTAACGCTTT 3030 3040 3050 3060 3070 TGCTCTANAT TTGGGAGGCT TANAGTGTTG TTTGCANTC CTACACGCGT GTCTANCTAG TGGAGGTTCG TCAGCTGCTC TAGTTTGAAT CATCATCGC GTAGTATTCC
ACGAGATTTA AACCCTCCGA ATTTCACAAC AAACGTTAGA GATGTCGCCA CAGATTGATC ACCTCCAAGC AGTCGACGAG ATCAAACTTA GTAGTAGCCG CATCATAAGG

3110	3120	3130	3140 * *	3150	3160	3170	3180	3190		3210
			ATTTCTCGTC	GAGAACGTTA	AAATAATCGT	TGTAACTCAC			TGTATTCTAC	
ATGAAAATGT	CAATCCTGTG	CCACATAACA	TAAAGAGCAG	CTCTTGCAAT	TTTATTAGCA	ACATTGAGTG	TAGGAAATAA	AATAGATATA	ACATAAGATG	AGGAAAGAAT
3220	2020	2244		22.50			2200	2200	2210	7770
* *	3230	3240	3250	3260	3270	3280 * *	3290	3300	3310	* *
									TAAATCATAT ATTTAGTATA	
INCOINMAI	MIGGCIIMII	CICINICOCI	ICCIIAAGAA	MANIMACIAN	TIGATCAGIT	INCICATALA	INTINACTIT	HOMITIMI	MITIAGIAIA	· · · · · · · · · · · · · · · · · · ·
3330	3340	3350	3360	3370	3380	3390	3400	3410	3420	3430
CGAAATATCA	GTAATAGACA								GCAAATACAG	CTTCATTCAA
									CGTTTATGTC	
3440	3450 * *				3490	3500	3510			3540
					TCAGATGATG	AGAAAGTAAA	TATAAATITA		TAATATAAT	
GCITAATGGA	MAMITAMAA	MGICIGIGIG	GARTARIGIT	TGATIGATIC	AGICIACIAC	TOTTICATTI	WIWIIIWWI	IGANIACCCA	ATTATATTA	TITCIAAGIA
3550	3560	3570	3580	3590	3600	3610	3620	3630	3640	3650
	* *	* *					* *	* *	* *	* *
									ATTAAAATAG TAATTTTATC	
3660	3670			3700	3710	3720	3730			3760
AGATGTAAAT	* *								GAATTTTTTA	
TCTACATTTA	TTAATAAACC	TCCATTTCCT	ATATTTTAAT	CAGATAGAAA	GTGTACCTTT	ACTTAATGGA	TTATAATTAT	TAATACTATC	CTTAAAAAAT	CCTAAATGTC
3770 * *	3780	3790 * *		3810						3870
									TAATAGCCAG	
GACAATATAC	MINGILGIIM	IGICCGICIA	GATACCAATA	CCATTTIGIG	ACATTGCCCT	TCGTCGTAAG	ATACCATTGA	CCGGATACAA	ATTATCGGTC	TAGTAAAATG
3880	3890	3900	3910	3920	3930	3940	3950	3960	3970	3980
* *	* *	* *	* *	* *		* *	* *	* *	* *	* *
									GAAGTATTTT CTTCATAAAA	
3990	4000	4010	4020	4030	4040	4050	4060		4080	4090
AGATAAAGAT	AGTCTATCTT	ATCTACAAGA	TATGAAAGAA	GATAATCATT	TAGTAGTAGC				CGTGGAAGCT	AATTATTTT
									GCACCTTCGA	
4100 * *	4110 * *	4120 * *	4130	4140 * *	4150	4160	4170	4180	4190 * *	4200 * *
									GTACATTAAT CATGTAATTA	
MIGGINIAN	IGNICITOIR	WILLIAM CONT.	CIGNATURIA	1101111010	MATTIMOGGI	ININGCIANG	MINIMANGIA	. GINIIGICAL	Cutatuuttu	GICACIAIAI
4210	4220	4230	4240	4250	4260	4270	4280	4290	4300	4310
* *		* *	* *		* *	* *	* *	* *	* *	* *
GACTTTGCTA	GATGTCTGAG	TTGATACGTT	CCTTATTCGT	TATACGGTTA	ATACAGATTA	TAAAATTGAA	ATCTTGATTT	TGCAAGATGG	AATACTAAAA TTATGATTTI	TATCCTATGC
4320	4330	4340	4350	4360	4370	4380	4390	4400	4410	4420
									TAAGTTTAAT	
									ATTCAAATTA	

4430 TTTATGAAGG TACC AAATACTTCC ATGG